

The affinity of the mutant for the column material is enhanced thereby, so that elution does not take place until the salt concentrations are high, that is to say in regions which are normally less contaminated by smaller foreign proteins.

### In the Claims

Please replace claims 37, 58, and 59 with the following claims re-written in clean form:

37. (Amended) The method as claimed in claim 35, wherein a peptide which has the sequence QAGTFALRGDNPQG (SEQ ID NO: 1) is inserted into the structural protein.

58. (Amended) The method as claimed in claim 27, wherein one or more insertions in VP3 is/are located before and/or after at least one amino acid in the sequence selected from the group consisting of YKQIS SQSGA (SEQ ID NO: 2), YLTLN NGSQA (SEQ ID NO: 3), YYLSR TNTPS (SEQ ID NO: 4), EEKFF PQSGV (SEQ ID NO: 5), NPVAT EQYGS (SEQ ID NOS: 6 and 7), LQRGN RQAAT (SEQ ID NO: 8), and NVDFT VDTNG (SEQ ID NO: 9)

59. (Amended) The method as claimed in claim 38, wherein one or more additional insertion(s) in VP3 is/are located before and/or after at least one amino acid in the sequence selected from the group consisting of YKQIS SQSGA (SEQ ID NO: 2), YLTLN NGSQA (SEQ ID NO: 3), YYLSR TNTPS (SEQ ID NO: 4), EEKFF PQSGV (SEQ ID NO: 5), NPVAT EQYGS (SEQ ID NOS: 6 and 7), LQRGN RQAAT (SEQ ID NO: 8), and NVDFT VDTNG (SEQ ID NO: 9).